

1600

15 DATE: 01/24/2003 DIS 2/11/03

TIME: 13:17:53

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/541,462B

Input Set: A:\5470-255.ST25.txt Output Set: N:\CRF4\01242003\I541462B.raw

```
3 <110> APPLICANT: University of North Carolina at Chapel Hill
             Xiong, Yue
             Ohta, Tomohiko
     7 <120> TITLE OF INVENTION: ISOLATION OF ROC1 AND ROC2
     9 <130> FILE REFERENCE: 5470-255
    11 <140> CURRENT APPLICATION NUMBER: US 09/541,462B
    12 <141> CURRENT FILING DATE: 2000-03-21
    14 <160> NUMBER OF SEQ ID NOS: 41
    16 <170> SOFTWARE: PatentIn version 3.1
    18 <210> SEQ ID NO: 1
    19 <211> LENGTH: 327
    20 <212> TYPE: DNA
    21 <213> ORGANISM: Homo sapiens
                                                                   ENTERED
    23 <220> FEATURE:
     24 <221> NAME/KEY: CDS
     25 <222> LOCATION: (1)..(324)
     26 <223> OTHER INFORMATION:
W--> 29 <400> 1
     30 atg gcg gca gcg atg gat gtg gat acc ccg agc ggc acc aac agc ggc
                                                                               48
     31 Met Ala Ala Ala Met Asp Val Asp Thr Pro Ser Gly Thr Asn Ser Gly
     34 gcg ggc aag aag cgc ttt gaa gtg aaa aag tgg aat gca gta gcc ctc
                                                                               96
     35 Ala Gly Lys Lys Arg Phe Glu Val Lys Lys Trp Asn Ala Val Ala Leu
                                        25
                                                                              144
     38 tgg gcc tgg gat att gtg gtt gat aac tgt gcc atc tgc agg aac cac
     39 Trp Ala Trp Asp Ile Val Val Asp Asn Cys Ala Ile Cys Arg Asn His
                35
                                    40
     40
                                                                              192
     42 att atg gat ctt tgc ata gaa tgt caa gct aac cag gcg tcc gct act
     43 Ile Met Asp Leu Cys Ile Glu Cys Gln Ala Asn Gln Ala Ser Ala Thr
                                55
     46 tca gaa gag tgt act gtc gca tgg gga gtc tgt aac cat gct ttt cac
                                                                              240
     47 Ser Glu Glu Cys Thr Val Ala Trp Gly Val Cys Asn His Ala Phe His
                                                75
                            70
     50 ttc cac tgc atc tct cgc tgg ctc aaa aca cga cag gtg tgt cca ttg
                                                                              288
     51 Phe His Cys Ile Ser Arg Trp Leu Lys Thr Arg Gln Val Cys Pro Leu
                                             90
                        85
                                                                              327
     54 gac aac aga gag tgg gaa ttc caa aag tat ggg cac tag
     55 Asp Asn Arg Glu Trp Glu Phe Gln Lys Tyr Gly His
     59 <210> SEQ ID NO: 2
     60 <211> LENGTH: 108
     61 <212> TYPE: PRT
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62 <213> ORGANISM: Homo sapiens

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64 <400> SEQUENCE: 2
    66 Met Ala Ala Ala Met Asp Val Asp Thr Pro Ser Gly Thr Asn Ser Gly
                                            10
    70 Ala Gly Lys Lys Arg Phe Glu Val Lys Lys Trp Asn Ala Val Ala Leu
                                        25
                    20
    71
    74 Trp Ala Trp Asp Ile Val Val Asp Asn Cys Ala Ile Cys Arg Asn His
    78 Ile Met Asp Leu Cys Ile Glu Cys Gln Ala Asn Gln Ala Ser Ala Thr
                                55
    82 Ser Glu Glu Cys Thr Val Ala Trp Gly Val Cys Asn His Ala Phe His
                                                75
                            70
    86 Phe His Cys Ile Ser Arg Trp Leu Lys Thr Arg Gln Val Cys Pro Leu
                        85
    90 Asp Asn Arg Glu Trp Glu Phe Gln Lys Tyr Gly His
                                        105
                   100
    94 <210> SEQ ID NO: 3
    95 <211> LENGTH: 342
     96 <212> TYPE: DNA
     97 <213> ORGANISM: Homo sapiens
     99 <220> FEATURE:
     100 <221> NAME/KEY: CDS
     101 <222> LOCATION: (1)..(339)
     102 <223> OTHER INFORMATION:
W--> 105 <400> 3
     106 atg gcc gac gtg gaa gac gga gag gaa acc tgc gcc ctg gcc tct cac
                                                                                48
     107 Met Ala Asp Val Glu Asp Gly Glu Glu Thr Cys Ala Leu Ala Ser His
                                                                  15
                                              10
     110 tee ggg age tea gge tea aeg teg gga gge gae aag atg tte tee ete
                                                                                96
     111 Ser Gly Ser Ser Gly Ser Thr Ser Gly Gly Asp Lys Met Phe Ser Leu
                                          25
                                                                                144
     114 aag aag tgg aac ceg gtg gcc atg tgg age tgg gac gtg gag tgc gat
     115 Lys Lys Trp Asn Pro Val Ala Met Trp Ser Trp Asp Val Glu Cys Asp
                                      40
     118 acg tgc gcc atc tgc agg gtc cag gtg atg gat gcc tgt ctt aga tgt
                                                                                192
     119 Thr Cys Ala Ile Cys Arg Val Gln Val Met Asp Ala Cys Leu Arg Cys
     120
             50
                                                                                240
     122 caa gct gaa aac aaa caa gag gac tgt gtt gtg gtc tgg gga gaa tgt
     123 Gln Ala Glu Asn Lys Gln Glu Asp Cys Val Val Val Trp Gly Glu Cys
                                                                                288
     126 aat cat too tto cac aac tgo tgo atg too ctg tgg gtg aaa cag aac
     127 Asn His Ser Phe His Asn Cys Cys Met Ser Leu Trp Val Lys Gln Asn
                         85
                                                                                336
     130 aat cgc tgc cct ctc tgc cag cag gac tgg gtg gtc caa aga atc ggc
     131 Asn Arg Cys Pro Leu Cys Gln Gln Asp Trp Val Val Gln Arg Ile Gly
                                                              110
                                          105
                     100
     132
                                                                                342
     134 aaa tga
     135 Lys
     139 <210> SEQ ID NO: 4
     140 <211> LENGTH: 113
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RAW SECUENCE LISTING

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141 <212> TYPE: PRT
142 <213> ORGANISM: Homo sapiens
144 <400> SEQUENCE: 4
146 Met Ala Asp Val Glu Asp Gly Glu Glu Thr Cys Ala Leu Ala Ser His
147 1
150 Ser Gly Ser Ser Gly Ser Thr Ser Gly Gly Asp Lys Met Phe Ser Leu
                                    25
151
154 Lys Lys Trp Asn Pro Val Ala Met Trp Ser Trp Asp Val Glu Cys Asp
          35
158 Thr Cys Ala Ile Cys Arg Val Gln Val Met Asp Ala Cys Leu Arg Cys
                            55
162 Gln Ala Glu Asn Lys Gln Glu Asp Cys Val Val Val Trp Gly Glu Cys
                        70
166 Asn His Ser Phe His Asn Cys Cys Met Ser Leu Trp Val Lys Gln Asn
                                        90
                   85
170 Asn Arg Cys Pro Leu Cys Gln Gln Asp Trp Val Val Gln Arg Ile Gly
               100
                                    105
171
174 Lys
178 <210> SEQ ID NO: 5
179 <211> LENGTH: 33
180 <212> TYPE: DNA
181 <213> ORGANISM: Artificial sequence
183 <220> FEATURE:
184 <223> OTHER INFORMATION: Synthetic oligonucleotide primer
186 <400> SEQUENCE: 5
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187 tttaaagaga aataggatcc catgagcaac gaa
190 <210> SEQ ID NO: 6
191 <211> LENGTH: 33
192 <212> TYPE: DNA
193 <213> ORGANISM: Artificial sequence
195 <220> FEATURE:
196 <223> OTHER INFORMATION: Synthetic oligonucleotide primer
198 <400> SEQUENCE: 6
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199 ttaaatgttt acggggaatt catttttca cct
202 <210> SEQ ID NO: 7
203 <211> LENGTH: 33
204 <212> TYPE: DNA
205 <213> ORGANISM: Artificial sequence
207 <220> FEATURE:
208 <223> OTHER INFORMATION: Synthetic oligonucleotide primer
210 <400> SEQUENCE: 7
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211 ggcaatacag attaggatcc tatgaaagtt aaa
214 <210> SEQ ID NO: 8
215 <211> LENGTH: 33
216 <212> TYPE: DNA
217 <213> ORGANISM: Artificial sequence
219 <220> FEATURE:
220 <223> OTHER INFORMATION: Synthetic oligonucleotide primer
222 <400> SEQUENCE: 8
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Input Set : A:\5470-255.ST25.txt

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223 aattgtgatt totagaatto tittitatog taa	33
226 <210> SEQ ID NO: 9	
227 <211> LENGTH: 33	
228 <212> TYPE: DNA	
229 <213> ORGANISM: Artificial sequence	
231 <220> FEATURE:	
232 <223> OTHER INFORMATION: Synthetic oligonucleotide primer	
234 <400> SEQUENCE: 9	33
235 atccccatgg ctatgataac taataagaaa ata	00
238 <210> SEQ 1D NO. 10	
239 <211> LENGIH: 33	
240 <212> TYPE: DNA	
241 <213> ORGANISM: Artificial sequence	
243 <220> FEATURE: 244 <223> OTHER INFORMATION: Synthetic oligonucleotide primer	
246 <400> SEQUENCE: 10	33
247 ctgcagagct cgttaggaaa ggtaatggta ata	
250 <210> SEQ ID NO: 11	
251 <211> LENGTH: 33	
252 <212> TYPE: DNA 253 <213> ORGANISM: Artificial sequence	
255 <220> FEATURE: 256 <223> OTHER INFORMATION: Synthetic oligonucleotide primer	
258 <400> SEQUENCE: 11	
258 44007 SEQUENCE: II 259 atcoccatgg ctatgataaa tgagagcgtt tcc	33
262 <210> SEQ ID NO: 12	
263 <211> LENGTH: 33	
264 <212> TYPE: DNA	
265 <213> ORGANISM: Artificial sequence	
267 <220> FEATURE:	
268 <223> OTHER INFORMATION: Synthetic oligonucleotide primer	
270 <400> SEQUENCE: 12	
271 agctcgtcga cattagtact tgtaagttgc tat	33
274 <210> SEQ ID NO: 13	
275 <211> LENGTH: 33	
276 <212> TYPE: DNA	
277 <213> ORGANISM: Artificial sequence	
279 <220> FEATURE:	
280 <223> OTHER INFORMATION: Synthetic oligonucleotide primer	
282 <400> SEQUENCE: 13	
283 atccccatgg ctatgtcatt tcagattacc cca	33
286 <210> SEQ ID NO: 14	
287 <211> LENGTH: 33	
288 <212> TYPE: DNA	
289 <213> ORGANISM: Artificial sequence	
291 <220> FEATURE:	
292 <223> OTHER INFORMATION: Synthetic oligonucleotide primer	
294 <400> SEQUENCE: 14	33
295 agctcgtcga catcatgagt ttttatgccc att	33

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Input Set : A:\5470-255.ST25.txt

Output Set: N:\CRF4\01242003\I541462B.raw

- 298 <210> SEQ ID NO: 15 299 <211> LENGTH: 15
- 300 <212> TYPE: PRT
- 301 <213> ORGANISM: Artificial sequence
- 303 <220> FEATURE:
- 304 <223> OTHER INFORMATION: Synthetic peptide
- 306 <400> SEQUENCE: 15
- 308 Cys Met Ala Ala Ala Met Asp Val Asp Thr Pro Ser Gly Thr Asn
- 309 1 5
- 312 <210> SEQ ID NO: 16
- 313 <211> LENGTH: 13 314 <212> TYPE: PRT
- 315 <213> ORGANISM: Artificial sequence
- 317 <220> FEATURE:
- 318 <223> OTHER INFORMATION: Synthetic peptide
- 320 <400> SEQUENCE: 16
- 322 Cys Asp Asn Arg Glu Trp Glu Phe Gln Lys Tyr Gly His
 - 23 1 5 1
- 326 <210> SEQ ID NO: 17
- 327 <211> LENGTH: 9
- 328 <212> TYPE: PRT
- 329 <213> ORGANISM: Artificial sequence
- 331 <220> FEATURE:
- 332 <223> OTHER INFORMATION: Synthetic peptide
- 334 <400> SEQUENCE: 17
- 336 Cys Arg Gln Glu Trp Lys Phe Lys Glu
- 337 1
- 340 <210> SEO ID NO: 18
- 341 <211> LENGTH: 14
- 342 <212> TYPE: PRT
- 343 <213> ORGANISM: Artificial sequence
- 345 <220> FEATURE:
- 346 <223> OTHER INFORMATION: Synthetic peptide
- 348 <400> SEQUENCE: 18
- 350 Cys Arg Ser Gln Ala Ser Ala Asp Glu Tyr Ser Tyr Val Ala 351 1 5 10
- 331 1
- 354 <210> SEQ ID NO: 19
- 355 <211> LENGTH: 60
- 356 <212> TYPE: DNA
- 357 <213> ORGANISM: Artificial sequence
- 359 <220> FEATURE:
- 360 <223> OTHER INFORMATION: Synthetic oligonucleotide primer
- 362 <400> SEQUENCE: 19
 363 ttctccagtg gcagagaact ttaaagagaa atagttcaac cggatccccg ggttaattaa
- 366 <210> SEQ ID NO: 20
- 367 <211> LENGTH: 60 368 <212> TYPE: DNA
- 369 <213> ORGANISM: Artificial sequence
- 371 <220> FEATURE:

60

VERIFICATION SUMMARY

PATENT APPLICATION: US/09/541,462B

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L:29 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1,Line#:26 L:105 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:3,Line#:102